



IFWO

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/820,155

DATE: 07/29/2004

TIME: 11:40:57

Input Set : A:\Sequence listing.ST25.txt  
 Output Set: N:\CRF4\07292004\J820155.raw

3 <110> APPLICANT: NatImmune A/S  
 4 Weilguny, Dietmar  
 5 Jensenius, Jens Christian  
 6 Kongerslev, Leif  
 7 Matthiesen, Finn  
 9 <120> TITLE OF INVENTION: Treatment of SARS in individuals  
 11 <130> FILE REFERENCE: P 774 US00  
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/820,155  
 14 <141> CURRENT FILING DATE: 2004-04-08  
 16 <160> NUMBER OF SEQ ID NOS: 137  
 18 <170> SOFTWARE: PatentIn version 3.1  
 20 <210> SEQ ID NO: 1  
 21 <211> LENGTH: 652  
 22 <212> TYPE: PRT  
 23 <213> ORGANISM: Homo sapiens  
 25 <400> SEQUENCE: 1  
 27 Met Ala Thr Ser Met Gly Leu Leu Leu Leu Leu Leu Leu Leu Leu Thr  
 28 1 5 10 15  
 31 Gln Pro Gly Ala Gly Thr Gly Ala Asp Thr Glu Ala Val Val Cys Val  
 32 20 25 30  
 35 Gly Thr Ala Cys Tyr Thr Ala His Ser Gly Lys Leu Ser Ala Ala Glu  
 36 35 40 45  
 39 Ala Gln Asn His Cys Asn Gln Asn Gly Gly Asn Leu Ala Thr Val Lys  
 40 50 55 60  
 43 Ser Lys Glu Glu Ala Gln His Val Gln Arg Val Leu Ala Gln Leu Leu  
 44 65 70 75 80  
 47 Arg Arg Glu Ala Ala Leu Thr Ala Arg Met Ser Lys Phe Trp Ile Gly  
 48 85 90 95  
 51 Leu Gln Arg Glu Lys Gly Lys Cys Leu Asp Pro Ser Leu Pro Leu Lys  
 52 100 105 110  
 55 Gly Phe Ser Trp Val Gly Gly Glu Asp Thr Pro Tyr Ser Asn Trp  
 56 115 120 125  
 59 His Lys Glu Leu Arg Asn Ser Cys Ile Ser Lys Arg Cys Val Ser Leu  
 60 130 135 140  
 63 Leu Leu Asp Leu Ser Gln Pro Leu Leu Pro Ser Arg Leu Pro Lys Trp  
 64 145 150 155 160  
 67 Ser Glu Gly Pro Cys Gly Ser Pro Gly Ser Pro Gly Ser Asn Ile Glu  
 68 165 170 175  
 71 Gly Phe Val Cys Lys Phe Ser Phe Lys Gly Met Cys Arg Pro Leu Ala  
 72 180 185 190  
 75 Leu Gly Gly Pro Gly Gln Val Thr Tyr Thr Thr Pro Phe Gln Thr Thr  
 76 195 200 205  
 79 Ser Ser Ser Leu Glu Ala Val Pro Phe Ala Ser Ala Ala Asn Val Ala

P,6

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80	210	215	220													
83	Cys	Gly	Glu	Gly	Asp	Lys	Glu	Thr	Gln	Ser	His	Tyr	Phe	Leu	Cys	
84	225		230				235							240		
87	Lys	Glu	Lys	Ala	Pro	Asp	Val	Phe	Asp	Trp	Gly	Ser	Ser	Gly	Pro	Leu
88							245			250				255		
91	Cys	Val	Ser	Pro	Lys	Tyr	Gly	Cys	Asn	Phe	Asn	Asn	Gly	Gly	Cys	His
92							260			265				270		
95	Gln	Asp	Cys	Phe	Glu	Gly	Gly	Asp	Gly	Ser	Phe	Leu	Cys	Gly	Cys	Arg
96							275			280				285		
99	Pro	Gly	Phe	Arg	Leu	Leu	Asp	Asp	Leu	Val	Thr	Cys	Ala	Ser	Arg	Asn
100							290			295				300		
103	Pro	Cys	Ser	Ser	Ser	Pro	Cys	Arg	Gly	Gly	Ala	Thr	Cys	Val	Leu	Gly
104	305						310				315				320	
107	Pro	His	Gly	Lys	Asn	Tyr	Thr	Cys	Arg	Cys	Pro	Gln	Gly	Tyr	Gln	Leu
108							325			330				335		
111	Asp	Ser	Ser	Gln	Leu	Asp	Cys	Val	Asp	Val	Asp	Glu	Cys	Gln	Asp	Ser
112							340			345				350		
115	Pro	Cys	Ala	Gln	Glu	Cys	Val	Asn	Thr	Pro	Gly	Gly	Phe	Arg	Cys	Glu
116							355			360				365		
119	Cys	Trp	Val	Gly	Tyr	Glu	Pro	Gly	Gly	Pro	Gly	Glu	Gly	Ala	Cys	Gln
120							370			375				380		
123	Asp	Val	Asp	Glu	Cys	Ala	Leu	Gly	Arg	Ser	Pro	Cys	Ala	Gln	Gly	Cys
124	385						385			390				395		400
127	Thr	Asn	Thr	Asp	Gly	Ser	Phe	His	Cys	Ser	Cys	Glu	Gly	Tyr	Val	
128							405			410				415		
131	Leu	Ala	Gly	Glu	Asp	Gly	Thr	Gln	Cys	Gln	Asp	Val	Asp	Glu	Cys	Val
132							420			425				430		
135	Gly	Pro	Gly	Gly	Pro	Leu	Cys	Asp	Ser	Leu	Cys	Phe	Asn	Thr	Gln	Gly
136							435			440				445		
139	Ser	Phe	His	Cys	Gly	Cys	Leu	Pro	Gly	Trp	Val	Leu	Ala	Pro	Asn	Gly
140							450			455				460		
143	Val	Ser	Cys	Thr	Met	Gly	Pro	Val	Ser	Leu	Gly	Pro	Pro	Ser	Gly	Pro
144	465						465			470				475		480
147	Pro	Asp	Glu	Glu	Asp	Lys	Gly	Glu	Lys	Glu	Gly	Ser	Thr	Val	Pro	Arg
148							485			490				495		
151	Ala	Ala	Thr	Ala	Ser	Pro	Thr	Arg	Gly	Pro	Glu	Gly	Thr	Pro	Lys	Ala
152							500			505				510		
155	Thr	Pro	Thr	Thr	Ser	Arg	Pro	Ser	Leu	Ser	Ser	Asp	Ala	Pro	Ile	Thr
156							515			520				525		
159	Ser	Ala	Pro	Leu	Lys	Met	Leu	Ala	Pro	Ser	Gly	Ser	Pro	Gly	Val	Trp
160							530			535				540		
163	Arg	Glu	Pro	Ser	Ile	His	His	Ala	Thr	Ala	Ala	Ser	Gly	Pro	Gln	Glu
164	545						545			550				555		560
167	Pro	Ala	Gly	Gly	Asp	Ser	Ser	Val	Ala	Thr	Gln	Asn	Asn	Asp	Gly	Thr
168							565			570				575		
171	Asp	Gly	Gln	Lys	Leu	Leu	Leu	Phe	Tyr	Ile	Leu	Gly	Thr	Val	Val	Ala
172							580			585				590		
175	Ile	Leu	Leu	Leu	Leu	Ala	Leu	Ala	Leu	Gly	Leu	Leu	Val	Tyr	Arg	Lys
176							595			600				605		

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179 Arg Arg Ala Lys Arg Glu Glu Lys Lys Glu Lys Pro Gln Asn Ala  
180 610 615 620  
183 Ala Asp Ser Tyr Ser Trp Val Pro Glu Arg Ala Glu Ser Arg Ala Met  
184 625 630 635 640  
187 Glu Asn Gln Tyr Ser Pro Thr Pro Gly Thr Asp Cys  
188 645 650  
191 <210> SEQ ID NO: 2  
192 <211> LENGTH: 742  
193 <212> TYPE: PRT  
194 <213> ORGANISM: Mus musculus  
196 <400> SEQUENCE: 2  
198 Met Lys Asp Asp Phe Ala Glu Glu Glu Val Gln Ser Phe Gly Tyr  
199 1 5 10 15  
202 Lys Arg Phe Gly Ile Gln Glu Gly Thr Gln Cys Thr Lys Cys Lys Asn  
203 20 25 30  
206 Asn Trp Ala Leu Lys Phe Ser Ile Val Leu Leu Tyr Ile Leu Cys Ala  
207 35 40 45  
210 Leu Leu Thr Ile Thr Val Ala Ile Leu Gly Tyr Lys Val Val Glu Lys  
211 50 55 60  
214 Met Asp Asn Val Thr Asp Gly Met Glu Thr Ser His Gln Thr Tyr Asp  
215 65 70 75 80  
218 Asn Lys Leu Thr Ala Val Glu Ser Asp Leu Lys Lys Leu Gly Asp Gln  
219 85 90 95  
222 Ala Gly Lys Lys Ala Leu Ser Thr Asn Ser Glu Leu Ser Thr Phe Arg  
223 100 105 110  
226 Ser Asp Ile Leu Asp Leu Arg Gln Gln Leu Gln Glu Ile Thr Glu Lys  
227 115 120 125  
230 Thr Ser Lys Asn Lys Asp Thr Leu Glu Lys Leu Gln Ala Asn Gly Asp  
231 130 135 140  
234 Ser Leu Val Asp Arg Gln Ser Gln Leu Lys Glu Thr Leu Gln Asn Asn  
235 145 150 155 160  
238 Ser Phe Leu Ile Thr Thr Val Asn Lys Thr Leu Gln Ala Tyr Asn Gly  
239 165 170 175  
242 Tyr Val Thr Asn Leu Gln Gln Asp Thr Ser Val Leu Gln Gly Asn Leu  
243 180 185 190  
246 Gln Ser Gln Met Tyr Ser Gln Ser Val Val Ile Met Asn Leu Asn Asn  
247 195 200 205  
250 Leu Asn Leu Thr Gln Val Gln Gln Arg Asn Leu Ile Ser Asn Leu Gln  
251 210 215 220  
254 Gln Ser Val Asp Asp Thr Ser Leu Ala Ile Gln Arg Ile Lys Asn Asp  
255 225 230 235 240  
258 Phe Gln Asn Leu Gln Gln Val Phe Leu Gln Ala Lys Lys Asp Thr Asp  
259 245 250 255  
262 Trp Leu Lys Glu Lys Val Gln Ser Leu Gln Thr Leu Ala Ala Asn Asn  
263 260 265 270  
266 Ser Ala Leu Ala Lys Ala Asn Asn Asp Thr Leu Glu Asp Met Asn Ser  
267 275 280 285  
270 Gln Leu Ser Ser Phe Thr Gly Gln Met Asp Asn Ile Thr Thr Ile Ser  
271 290 295 300

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274 Gln Ala Asn Glu Gln Ser Leu Lys Asp Leu Gln Asp Leu His Lys Asp  
 275 305 310 315 320  
 278 Thr Glu Asn Arg Thr Ala Val Lys Phe Ser Gln Leu Glu Glu Arg Phe  
 279 325 330 335  
 282 Gln Val Phe Glu Thr Asp Ile Val Asn Ile Ile Ser Asn Ile Ser Tyr  
 283 340 345 350  
 286 Thr Ala His His Leu Arg Thr Leu Thr Ser Asn Leu Asn Asp Val Arg  
 287 355 360 365  
 290 Thr Thr Cys Thr Asp Thr Leu Thr Arg His Thr Asp Asp Leu Thr Ser  
 291 370 375 380  
 294 Leu Asn Asn Thr Leu Val Asn Ile Arg Leu Asp Ser Ile Ser Leu Arg  
 295 385 390 395 400  
 298 Met Gln Gln Asp Met Met Arg Ser Lys Leu Asp Thr Glu Val Ala Asn  
 299 405 410 415  
 302 Leu Ser Val Val Met Glu Glu Met Lys Leu Val Asp Ser Lys His Gly  
 303 420 425 430  
 306 Gln Leu Ile Lys Asn Phe Thr Ile Leu Gln Gly Pro Pro Gly Pro Arg  
 307 435 440 445  
 310 Gly Pro Lys Gly Asp Arg Gly Ser Gln Gly Pro Pro Gly Pro Thr Gly  
 311 450 455 460  
 314 Asn Lys Gly Gln Lys Gly Glu Lys Gly Glu Pro Gly Pro Pro Gly Pro  
 315 465 470 475 480  
 318 Ala Gly Glu Arg Gly Thr Ile Gly Pro Val Gly Pro Pro Gly Glu Arg  
 319 485 490 495  
 322 Gly Ser Lys Gly Ser Lys Gly Ser Gln Gly Pro Lys Gly Ser Arg Gly  
 323 500 505 510  
 326 Ser Pro Gly Lys Pro Gly Pro Gln Gly Pro Ser Gly Asp Pro Gly Pro  
 327 515 520 525  
 330 Pro Gly Pro Pro Gly Lys Asp Gly Leu Pro Gly Pro Gln Gly Pro Pro  
 331 530 535 540  
 334 Gly Phe Gln Gly Leu Gln Gly Thr Val Gly Glu Pro Gly Val Pro Gly  
 335 545 550 555 560  
 338 Pro Arg Gly Leu Pro Gly Leu Pro Gly Val Pro Gly Met Pro Gly Pro  
 339 565 570 575  
 342 Lys Gly Pro Pro Gly Pro Pro Gly Pro Ser Gly Ala Met Glu Pro Leu  
 343 580 585 590  
 346 Ala Leu Gln Asn Glu Pro Thr Pro Ala Ser Glu Val Asn Gly Cys Pro  
 347 595 600 605  
 350 Pro His Trp Lys Asn Phe Thr Asp Lys Cys Tyr Tyr Phe Ser Leu Glu  
 351 610 615 620  
 354 Lys Glu Ile Phe Glu Asp Ala' Lys Leu Phe Cys Glu Asp Lys Ser Ser  
 355 625 630 635 640  
 358 His Leu Val Phe Ile Asn Ser Arg Glu Glu Gln Gln Trp Ile Lys Lys  
 359 645 650 655  
 362 His Thr Val Gly Arg Glu Ser His Trp Ile Gly Leu Thr Asp Ser Glu  
 363 660 665 670  
 366 Gln Glu Ser Glu Trp Lys Trp Leu Asp Gly Ser Pro Val Asp Tyr Lys  
 367 675 680 685  
 370 Asn Trp Lys Ala Gly Gln Pro Asp Asn Trp Gly Ser Gly His Gly Pro

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371 690 695 700  
 374 Gly Glu Asp Cys Ala Gly Leu Ile Tyr Ala Gly Gln Trp Asn Asp Phe  
 375 705 710 715 720  
 378 Gln Cys Asp Glu Ile Asn Asn Phe Ile Cys Glu Lys Glu Arg Glu Ala  
 379 725 730 735  
 382 Val Pro Ser Ser Ile Leu  
 383 740  
 386 <210> SEQ ID NO: 3  
 387 <211> LENGTH: 371  
 388 <212> TYPE: PRT  
 389 <213> ORGANISM: Bos taurus  
 391 <400> SEQUENCE: 3  
 393 Met Leu Leu Leu Pro Leu Ser Val Leu Leu Leu Thr Gln Pro Trp  
 394 1 5 10 15  
 397 Arg Ser Leu Gly Ala Glu Met Lys Ile Tyr Ser Gln Lys Thr Leu Ala  
 398 20 25 30  
 401 Asn Gly Thr Leu Val Val Cys Arg Pro Pro Glu Gly Gly Leu Pro  
 402 35 40 45  
 405 Gly Arg Asp Gly Gln Asp Gly Arg Glu Gly Pro Gln Gly Glu Lys Gly  
 406 50 55 60  
 409 Asp Pro Gly Ser Pro Gly Pro Ala Gly Arg Ala Gly Arg Pro Gly Pro  
 410 65 70 75 80  
 413 Ala Gly Pro Ile Gly Pro Lys Gly Asp Asn Gly Ser Ala Gly Glu Pro  
 414 85 90 95  
 417 Gly Pro Lys Gly Asp Thr Gly Pro Pro Gly Pro Pro Gly Met Pro Gly  
 418 100 105 110  
 421 Pro Ala Gly Arg Glu Gly Pro Ser Gly Lys Gln Gly Ser Met Gly Pro  
 422 115 120 125  
 425 Pro Gly Thr Pro Gly Pro Lys Gly Asp Thr Gly Pro Lys Gly Met  
 426 130 135 140  
 429 Gly Ala Pro Gly Met Gln Gly Ser Pro Gly Pro Ala Gly Leu Lys Gly  
 430 145 150 155 160  
 433 Glu Arg Gly Ala Pro Gly Glu Leu Gly Ala Pro Gly Ser Ala Gly Val  
 434 165 170 175  
 437 Ala Gly Pro Ala Gly Ala Ile Gly Pro Gln Gly Pro Ser Gly Ala Arg  
 438 180 185 190  
 441 Gly Pro Pro Gly Leu Lys Gly Asp Arg Gly Asp Pro Gly Glu Arg Gly  
 442 195 200 205  
 445 Ala Lys Gly Glu Ser Gly Leu Ala Asp Val Asn Ala Leu Lys Gln Arg  
 446 210 215 220  
 449 Val Thr Ile Leu Glu Gly Gln Leu Gln Arg Leu Gln Asn Ala Phe Ser  
 450 225 230 235 240  
 453 Arg Tyr Lys Lys Ala Val Leu Phe Pro Asp Gly Gln Ala Val Gly Lys  
 454 245 250 255  
 457 Lys Ile Phe Lys Thr Ala Gly Ala Val Lys Ser Tyr Ser Asp Ala Gln  
 458 260 265 270  
 461 Gln Leu Cys Arg Glu Ala Lys Gly Gln Leu Ala Ser Pro Arg Ser Ala  
 462 275 280 285  
 465 Ala Glu Asn Glu Ala Val Ala Gln Leu Val Arg Ala Lys Asn Asn Asp

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 07/29/2004  
PATENT APPLICATION: US/10/820,155 TIME: 11:40:59

Input Set : A:\Sequence listing.ST25.txt  
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**Please Note:**

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:37; Xaa Pos. 7  
Seq#:75; Xaa Pos. 7  
Seq#:94; Xaa Pos. 91,92,93,94,95,96,97,98,99,100,101,102,103,104,105,106  
Seq#:94; Xaa Pos. 107,108,109,110,111,112,113,114,115,116

**VERIFICATION SUMMARY**

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Input Set : A:\Sequence listing.ST25.txt  
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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:4011 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0  
L:9259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:75 after pos.:0  
L:11454 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:80  
L:11458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:96  
L:11462 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:112